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Phe	Thr	Phe	Thr	Trp	Ala	Phe	Gln	Arg	Thr	Asn	Gln	Gly	Gln	Asp	Asn
				565				570				575			
Arg	Arg	Phe	Ile	Asn	Asp	Met	Val	Lys	Ile	Tyr	Ser	Ile	Thr	Ala	Thr
				580				585				590			
Asn	Ala	Val	Asp	Gly	Val	Ala	Ser	Ser	Cys	Arg	Ala	Cys	Ala	Leu	Gly
				595				600				605			
Ser	Glu	Gln	Ser	Gly	Ser	Ser	Cys	Val	Pro	Cys	Pro	Pro	Gly	His	Tyr
				610				615				620			
Ile	Glu	Lys	Glu	Thr	Asn	Gln	Cys	Lys	Glu	Cys	Pro	Pro	Asp	Thr	Tyr
				625				630				635			
Leu	Ser	Ile	His	Gln	Val	Tyr	Gly	Lys	Glu	Ala	Cys	Ile	Pro	Cys	Gly
				645				650				655			
Pro	Gly	Ser	Lys	Asn	Asn	Gln	Asp	His	Ser	Val	Cys	Tyr	Ser	Asp	Cys
				660				665				670			
Phe	Phe	Tyr	His	Glu	Lys	Glu	Asn	Gln	Ile	Leu	His	Tyr	Asp	Phe	Ser
				675				680				685			

Asn	Leu	Ser	Ser	Val	Gly	Ser	Leu	Met	Asn	Gly	Pro	Ser	Phe	Thr	Ser		
690						695					700						
Lys	Gly	Thr	Lys	Tyr	Phe	His	Phe	Phe	Asn	Ile	Ser	Leu	Cys	Gly	His		
705					710					715					720		
Glu	Gly	Lys	Lys	Met	Ala	Leu	Cys	Thr	Asn	Asn	Ile	Thr	Asp	Phe	Thr		
				725					730					735			
Val	Lys	Glu	Ile	Val	Ala	Gly	Ser	Asp	Asp	Tyr	Thr	Asn	Leu	Val	Gly		
			740					745					750				
Ala	Phe	Val	Cys	Gln	Ser	Thr	Ile	Ile	Pro	Ser	Glu	Ser	Lys	Gly	Phe		
		755					760					765					
Arg	Ala	Ala	Leu	Ser	Ser	Gln	Ser	Ile	Ile	Leu	Ala	Asp	Thr	Phe	Ile		
770						775					780						
Gly	Val	Thr	Val	Glu	Thr	Thr	Leu	Lys	Asn	Ile	Asn	Ile	Lys	Glu	Asp		
785					790					795					800		
Met	Phe	Pro	Val	Pro	Thr	Ser	Gln	Ile	Pro	Asp	Val	His	Phe	Phe	Tyr		
				805					810					815			
Lys	Ser	Ser	Thr	Ala	Thr	Thr	Ser	Cys	Ile	Asn	Gly	Arg	Ser	Thr	Ala		
			820					825					830				
Val	Lys	Met	Arg	Cys	Asn	Pro	Thr	Lys	Ser	Gly	Ala	Gly	Val	Ile	Ser		
		835					840					845					
Val	Pro	Ser	Lys	Cys	Pro	Ala	Gly	Thr	Cys	Asp	Gly	Cys	Thr	Phe	Tyr		
	850					855					860						
Phe	Leu	Trp	Glu	Ser	Ala	Glu	Ala	Cys	Pro	Leu	Cys	Thr	Glu	His	Asp		
865					870					875					880		
Phe	His	Glu	Ile	Glu	Gly	Ala	Cys	Lys	Arg	Gly	Phe	Gln	Glu	Thr	Leu		
				885					890					895			
Tyr	Val	Trp	Asn	Glu	Pro	Lys	Trp	Cys	Ile	Lys	Gly	Ile	Ser	Leu	Pro		
			900					905					910				
Glu	Lys	Lys	Leu	Ala	Thr	Cys	Glu	Thr	Val	Asp	Phe	Trp	Leu	Lys	Val		
		915					920					925					
Gly	Ala	Gly	Val	Gly	Ala	Phe	Thr	Ala	Val	Leu	Leu	Val	Ala	Leu	Thr		
	930					935					940						
Cys	Tyr	Phe	Trp	Lys	Lys	Asn	Gln	Lys	Leu	Glu	Tyr	Lys	Tyr	Ser	Lys		
945					950					955					960		
Leu	Val	Met	Thr	Thr	Asn	Ser	Lys	Glu	Cys	Glu	Leu	Pro	Ala	Ala	Asp		
				965					970					975			
Ser	Cys	Ala	Ile	Met	Glu	Gly	Glu	Asp	Asn	Glu	Glu	Glu	Val	Val	Tyr		
			980					985					990				
Ser	Asn	Lys	Gln	Ser	Leu	Leu	Gly	Lys	Leu	Lys	Ser	Leu	Ala	Thr	Lys		
		995					1000						1005				

Glu Lys Glu Asp His Phe Glu Ser Val Gln Leu Lys Thr Ser Arg
 1010 1015 1020

Ser Pro Asn Ile
 1025

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<400> 5

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 1 5 10 15

Thr Leu Ala Thr Ala Asp Ile Pro Thr Ser Ser Leu Pro His Ala Pro
 20 25 30

Val Asn Gly Ala Cys Asp Glu Gly Glu Tyr Leu Asp Lys Arg His Asn
 35 40 45

Gln Cys Cys Asn Gln Cys Pro Pro Gly Glu Phe Ala Lys Val Arg Cys
 50 55 60

Asn Gly Asn Asp Asn Thr Lys Cys Glu Arg Cys Pro Pro His Thr Tyr
 65 70 75 80

Thr Ala Ile Pro Asn Tyr Ser Asn Gly Cys His Gln Cys Arg Lys Cys
 85 90 95

Pro Thr Gly Ser Phe Asp Lys Val Lys Cys Thr Gly Thr Gln Asn Ser
 100 105 110

Lys Cys Ser Cys Leu Pro Gly Trp Tyr Cys Ala Thr Asp Ser Ser Gln
 115 120 125

Thr Glu Asp Cys Arg Asp Cys Ile Pro Lys Arg Arg Cys Pro Cys Gly
 130 135 140

Tyr Phe Gly Gly Ile Asp Glu Gln Gly Asn Pro Ile Cys Lys Ser Cys
 145 150 155 160

Cys Val Gly Glu Tyr Cys Asp Tyr Leu Arg Asn Tyr Arg Leu Asp Pro
 165 170 175

Phe Pro Pro Cys Lys Leu Ser Lys Cys Asn
 180 185

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 <212> PRT
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<400> 6

Met Cys Val Gly Ala Arg Arg Leu Gly Arg Gly Pro Cys Ala Ala Leu
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Leu Leu Leu Gly Leu Gly Leu Ser Thr Val Thr Gly Leu His Cys Val
 20 25 30

Gly Asp Thr Tyr Pro Ser Asn Asp Arg Cys Cys His Glu Cys Arg Pro
35 40 45

Gly Asn Gly Met Val Ser Arg Cys Ser Arg Ser Gln Asn Thr Val Cys
50 55 60

Arg Pro Cys Gly Pro Gly Phe Tyr Asn Asp Val Val Ser Ser Lys Pro
65 70 75 80

Cys Lys Pro Cys Thr Trp Cys Asn Leu Arg Ser Gly Ser Glu Arg Lys
85 90 95

Gln Leu Cys Thr Ala Thr Gln Asp Thr Val Cys Arg Cys Arg Ala Gly
100 105 110

Thr Gln Pro Leu Asp Ser Tyr Lys Pro Gly Val Asp Cys Ala Pro Cys
115 120 125

Pro Pro Gly His Phe Ser Pro Gly Asp Asn Gln Ala Cys Lys Pro Trp
130 135 140

Thr Asn Cys Thr Leu Ala Gly Lys His Thr Leu Gln Pro Ala Ser Asn
145 150 155 160

Ser Ser Asp Ala Ile Cys Glu Asp Arg Asp Pro Pro Ala Thr Gln Pro
165 170 175

Gln Glu Thr Gln Gly Pro Pro Ala Arg Pro Ile Thr Val Gln Pro Thr
180 185 190

Glu Ala Trp Pro Arg Thr Ser Gln Gly Pro Ser Thr Arg Pro Val Glu
195 200 205

Val Pro Gly Gly Arg Ala Val Ala Ala Ile Leu Gly Leu Gly Leu Val
210 215 220

Leu Gly Leu Leu Gly Pro Leu Ala Ile Leu Leu Ala Leu Tyr Leu Leu
225 230 235 240

Arg Arg Asp Gln Arg Leu Pro Pro Asp Ala His Lys Pro Pro Gly Gly
245 250 255

Gly Ser Phe Arg Thr Pro Ile Gln Glu Glu Gln Ala Asp Ala His Ser
260 265 270

Thr Leu Ala Lys Ile
275

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Pro Cys Gln Glu Lys Asp Tyr His
1 5

<210> 8
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Gly Lys Glu Cys Thr Phe Ser Cys
1 5

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Gly Cys Asn Asn Ser Ser Trp Ile
1 5

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Phe Glu Phe Phe Ile Gln Asn Asp
1 5

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Gly Ser His Ser Val Met Leu Lys
1 5

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Thr Ile Glu Gly Val Ala Tyr Thr
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Ser Gln Phe Ser Gly Ser Ser Glu
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Glu Glu Gly Lys Thr Gln Ile Met
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Asp Gly Thr Lys Glu Cys Arg Pro
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Asp Gly Met Asn Gly Trp Glu Val
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Pro Gly Phe Lys Pro Pro Thr Ser
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Tyr Phe Met Val Asp Ile Asn Arg
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Gln Cys Gln Asp Asn Arg Arg Phe
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Lys Asn Asn Gln Asp His Ser Val
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<210> 21

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<400> 21

Cys Gly His Glu Gly Lys Lys Met
1 5

<210> 22

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<400> 22

Asp Thr Phe Ile Gly Val Thr Val
1 5

<210> 23

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<212> PRT

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<400> 23

Phe Phe Tyr Lys Ser Ser Thr Ala
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<210> 24

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<400> 24

Ile Ser Val Pro Ser Lys Cys Pro
1 5

<210> 25

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<400> 25

Arg Gly Phe Gln Glu Thr Leu Tyr
1 5

<210> 26

<211> 8

<212> PRT

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<400> 26

Lys Asn Gln Lys Lys Lys Lys Thr
1 5

<210> 27

<211> 8

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<213> human

<400> 27

Lys Asn Gln Lys Leu Glu Tyr Lys
1 5

<210> 28

<211> 8

<212> PRT

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<400> 28

Leu Ala Thr Lys Glu Lys Glu Asp
1 5

<210> 29

<211> 43

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<400> 29

Met Ala Pro Trp Asn Val Leu Pro Gly Pro His Phe Pro His Ser Ser
1 5 10 15

Arg Leu His Gly Ser Gly His Ser Arg Leu Ala Ala Ala Ala Ile Ser
20 25 30

Ile Ala Leu Lys Ala Phe Ser Cys Ala Ser Gly
35 40

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<211> 9

<212> PRT

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<400> 30

Thr Ile Glu Glu Glu Gly Ser Ser Glu
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<210> 31

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<400> 31

Cys Thr Glu Arg Pro Pro Cys Thr Thr Lys Asp Tyr Phe Gln Ile His
1 5 10 15

Thr Pro Cys Asp Glu Glu Gly Lys Thr Gln Ile Met Tyr Lys Trp Ile
20 25 30

Glu Pro Lys Ile Cys Arg Glu Asp Leu Thr Asp Ala Ile Arg Leu Pro
35 40 45

Pro Ser Gly Glu Lys Lys Asp Cys Pro Pro Cys Asn Pro Gly Phe Tyr
50 55 60

Asn Asn Gly Ser Ser Ser Cys His Pro Cys
65 70

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<400> 32

Thr Lys Gly Trp Trp Ile Ile Ser Gly Ser Ser Ser Leu Arg Arg Thr
1 5 10 15

Phe Lys His Ala Phe Cys Ser Thr Phe Ala Ala Glu Cys
20 25

<210> 33
<211> 35
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<400> 33

Phe Lys Met Asp Gly Ile Ile Tyr Ser Lys Arg Phe Lys His Ile Thr
1 5 10 15

Ile Val Met Trp Thr Gln Cys Leu Gln Arg Val Trp Thr Gly Met Ile
20 25 30

Lys Pro Pro
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<210> 34
<211> 37
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<400> 34

Gln Asp Asn Arg Pro Ile Pro Pro Leu Ser Ile Ser Ile Val Pro Tyr
1 5 10 15

Val Ser Ile Val Ala Gly Leu Ile Leu Trp Ile Ser Ile Asp Val Thr
20 25 30

Phe Pro Arg Arg Phe
35

<210> 35
<211> 78
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<400> 35

Lys Asn Gln Lys Leu Glu Tyr Lys Tyr Ser Lys Leu Val Met Thr Thr
1 5 10 15

Asn Ser Lys Glu Cys Glu Leu Pro Ala Ala Asp Ser Cys Ala Ile Met
20 25 30

Glu Gly Glu Asp Asn Glu Glu Glu Val Val Tyr Ser Asn Lys Gln Ser
35 40 45

Leu Leu Gly Lys Leu Lys Ser Leu Ala Thr Lys Glu Lys Glu Asp His
50 55 60

Phe Glu Ser Val Gln Leu Lys Thr Ser Arg Ser Pro Asn Ile
65 70 75

<210> 36

<211> 46

<212> DNA

<213> human

<400> 36

gcagcacata tgggggacct gccctcctcc tccagccgcc cgcttc

46

<210> 37

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<212> DNA

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<400> 37

gcagcaacta gtttagtcaa ccgtttcaca ggttgccaac tttttc

46

<210> 38

<211> 42

<212> DNA

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<400> 38

gcagcaggta cctcatatat ttgggggatct tgaggttttc ag

42

<210> 39

<211> 48

<212> DNA

<213> human

<400> 39

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48

<210> 40

<211> 27

<212> DNA

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<400> 40

gcagcacata tgctgttccg cgccccgg

27

<210> 41

<211> 59

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<212> DNA
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<400> 41
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<210> 42
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<212> DNA
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<400> 42
gggatccgga gcccaaattct tctgacaaaa ctcacacatg cccaccgtgc ccagcacctg 60
aattcgaggg tgcaccgtca gtcttctctt tcccccaaaa acccaaggac accctcatga 120
tctcccgga tcttgagggt acatgcgtgg tgggtggacgt aagccacgaa gaccctgagg 180
tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240
aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300
ggctgaatgg caaggagtac aagtgcagg tctccaacaa agccttccca acccccatcg 360
agaaaacat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc 420
catcccgga tgagctgacc aagaaccagg tcagcctgac ctgcctgggc aaaggcttct 480
atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga 540
ccacgcctcc cgtgctggac tccgacggct cttcttctct ctacagcaag ctcaccgtgg 600
acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggtctctg 660
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gactctagag gat 733

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<400> 43
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Gly Ser Gly Ile Lys Phe Asp Glu Trp Asp Glu Leu Pro Ala Gly Phe
20 25 30
Ser Asn Ile Ala Thr Phe Met Asp Thr Val Val Gly Pro Ser Asp Ser
35 40 45
Arg Pro Asp Gly Cys Asn Asn Ser Ser Trp Ile Pro Arg Gly Asn Tyr
50 55 60
Ile Glu Ser Asn Arg Asp Asp Cys Thr Val Ser Leu Ile Tyr Ala Val
65 70 75 80
His Leu Lys Lys Ser Gly Tyr Val Phe Phe Glu Tyr Gln Tyr Val Asp

85

90

95

Asn Asn Ile Phe Phe Glu Phe Phe Ile Gln Asn Asp Gln Cys Gln Glu
 100 105 110
 Met Asp Thr Thr Thr Asp Lys Trp Val Lys Leu Thr Asp Asn Gly Glu
 115 120 125
 Trp Gly Ser His Ser Val Met Leu Lys Ser Gly Thr Asn Ile Leu Tyr
 130 135 140
 Trp Arg Thr Thr Gly Ile Leu Met Gly Ser Lys Ala Val Lys Pro Val
 145 150 155 160
 Leu Val Lys Asn Ile Thr Ile Glu Gly Val Ala Tyr Thr Ser Glu Cys
 165 170 175
 Phe Pro Cys Lys Pro Gly Thr Phe Ser Asn Lys Pro Gly Ser Phe Asn
 180 185 190
 Cys Gln Val Cys Pro Arg Asn Thr Tyr Ser Glu Lys Gly Ala Lys Glu
 195 200 205
 Cys Ile Arg Cys Lys Asp Asp Ser Gln Phe Ser Glu Glu Gly Ser Ser
 210 215 220
 Glu Cys Thr Glu Arg Pro Pro Cys Thr Thr Lys Asp Tyr Phe Gln Ile
 225 230 235 240
 His Thr Pro Cys Asp Glu Glu Gly Lys Thr Gln Ile Met Tyr Lys Trp
 245 250 255
 Ile Glu Pro Lys Ile Cys Arg Glu Asp Leu Thr Asp Ala Ile Arg Leu
 260 265 270
 Pro Pro Ser Gly Glu Lys Lys Asp Cys Pro Pro Cys Asn Pro Gly Phe
 275 280 285
 Tyr Asn Asn Gly Ser Ser Ser Cys His Pro Cys Pro Pro Gly Thr Phe
 290 295 300
 Ser Asp Gly Thr Lys Glu Cys Arg Pro Cys Pro Ala Gly Thr Glu Pro
 305 310 315 320
 Ala Leu Gly Phe Glu Tyr Lys Trp Trp Asn Val Leu Pro Gly Asn Met
 325 330 335
 Lys Thr Ser Cys Phe Asn Val Gly Asn Ser Lys Cys Asp Gly Met Asn
 340 345 350
 Gly Trp Glu Val Ala Gly Asp His Ile Gln Ser Gly Ala Gly Gly Ser
 355 360 365
 Asp Asn Asp Tyr Leu Ile Leu Asn Leu His Ile Pro Gly Phe Lys Pro
 370 375 380
 Pro Thr Ser Met Thr Gly Ala Thr Gly Ser Glu Leu Gly Arg Ile Thr
 385 390 395 400
 Phe Val Phe Glu Thr Leu Cys Ser Ala Asp Cys Val Leu Tyr Phe Met

405

410

415

Val Asp Ile Asn Arg Lys Ser Thr Asn Val Val Glu Ser Trp Gly Gly
420 425 430

Thr Lys Glu Lys Gln Ala Tyr Thr His Ile Ile Phe Lys Asn Ala Thr
435 440 445

Phe Thr Phe Thr Trp Gly Ile Pro Arg Glu Leu Ile Gln Gly Pro Arg
450 455 460

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